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     Fungal metabolite screening: database of 474 mycotoxins and fungal
TI
     metabolites for dereplication by standardised liquid chromatography-UV-
     mass spectrometry methodology
ΑU
     Nielsen, Kristian Fog; Smedsgaard, Jorn
CS
     BioCentrum-DTU, Mycology Group, Technical University of Denmark, Lyngby,
     DK-2800, Den.
SO
     Journal of Chromatography, A (2003), 1002(1-2), 111-136
     CODEN: JCRAEY; ISSN: 0021-9673
PΒ
     Elsevier Science B.V.
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AN
     2003:279227 CAPLUS
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     138:270330
     Natural products: Discovery and screening
ΤI
AU
     Hilton, Matthew D.
     Bioprocess Fermentation Development, Eli Lilly and Company, Indianapolis,
CS
SO
     Handbook of Industrial Cell Culture (2003), 107-136.
                                                            Editor(s): Vinci,
     Victor A.; Parekh, Sarad R. Publisher: Humana Press Inc., Totowa, N. J.
     CODEN: 69DSXR; ISBN: 1-58829-032-8
DT
     Conference; General Review
     English
LA
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82

- L1ANSWER 3 OF 12 CAPLUS COPYRIGHT 2004 ACS on STN 2002:289965 CAPLUS AN DN 137:163136 TIElectrospray mass spectrometry applications in combinatorial chemistry ΑU Lee, Mike S. CS Milestone Development Services, Newtown, PA, USA SO Practical Spectroscopy (2002), 32 (Applied Electrospray Mass Spectrometry), 187-210 CODEN: PSPED9; ISSN: 0148-9054 PΒ Marcel Dekker, Inc. DT Journal; General Review LA English RE.CNT 63 THERE ARE 63 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT L1ANSWER 4 OF 12 CAPLUS COPYRIGHT 2004 ACS on STN 2002:30631 CAPLUS AN DN 136:315140 Automated molecular weight assignment of electrospray ionization mass TIspectra Williams, Jon D.; Weiner, Brian E.; Ormand, James R.; Brunner, Jimmy; ΑU Thornquest, Alan D., Jr.; Burinsky, David J. Preclinical Development Division, GlaxoSmithKline, Research Triangle Park, CS NC, 27709-3398, USA SO Rapid Communications in Mass Spectrometry (2001), 15(24), 2446-2455 CODEN: RCMSEF; ISSN: 0951-4198 PB John Wiley & Sons Ltd. DTJournal LA English RE.CNT 16 THERE ARE 16 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT L1ANSWER 5 OF 12 CAPLUS COPYRIGHT 2004 ACS on STN ΑN 2001:880037 CAPLUS DN 136:98264 TIMembrane-associated quinoprotein formaldehyde dehydrogenase from Methylococcus capsulatus Bath Zahn, James A.; Bergmann, David J.; Boyd, Jeffery M.; Kunz, Ryan C.; ΑU DiSpirito, Alan A. Department of Microbiology, Iowa State University, Ames, IA, 50011, USA CS Journal of Bacteriology (2001), 183(23), 6832-6840 SO CODEN: JOBAAY; ISSN: 0021-9193 PBAmerican Society for Microbiology DTJournal LΑ English RE.CNT 56 THERE ARE 56 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT T.7 ANSWER 6 OF 12 CAPLUS COPYRIGHT 2004 ACS on STN AN 2001:835758 CAPLUS DN 136:155824 TIOrganic pollutants in aqueous-solid phase environments: Types, analyses and characterizations AU Aboul-Kassim, Tarek A. T.; Simoneit, Bernd R. T. CS Department of Civil, Construction and Environmental Engineering, College of Engineering, Oregon State University, Corvallis, OR, 97331, USA
- DT Conference; General Review LA English

CODEN: 45NZAP

Springer, Berlin, Germany.

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     2001:381613 CAPLUS
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     Fast, efficient separations in drug discovery - LC-MS analysis using
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     column switching and rapid gradients
ΑU
     Needham, Shane R.; Wehr, Tim
CS
     Alturas Analytics Inc., Moscow, ID, USA
SO
     LC-GC Europe (2001), 14(4), 244,246,248-249
     CODEN: LCGCB4
PΒ
     Advanstar Communications, Inc.
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     Use of direct-infusion electrospray mass spectrometry to quide empirical
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     development of improved conditions for expression of secondary metabolites
     from actinomycetes
ΑU
     Zahn, James A.; Higgs, Richard E.; Hilton, Matthew D.
CS
     Natural Products Research, Eli Lilly and Company, Indianapolis, IN, 46285,
     USA
SO
     Applied and Environmental Microbiology (2001), 67(1), 377-386
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     Natural Products Research, Lilly Corporate Center, Eli Lilly and Company,
CS
     Indianapolis, IN, 46285, USA
SO
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     Directions in discovery: Fast, efficient separations in drug
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     discovery-LC-MS analysis using column switching and rapid gradients
AU
     Needham, S. R.
CS
     Alturas Analytics, Inc., Alturas Technology Park, Moscow, 83843, Russia
SO
     LC-GC (2000), 18(11), 1156, 1158, 1160-1161
     CODEN: LCGCE7; ISSN: 0888-9090
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      131:266391
      The role of mass spectrometry in the drug discovery process
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ΑU
      Pramanik, Birendra N.; Bartner, Peter L.; Chen, Guodong
 CS
      Schering-Plough Research Institute, Kenilworth, NJ, 07033, USA
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      Current Opinion in Drug Discovery & Development (1999), 2(4), 401-417
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     High-performance liquid chromatographic-electrospray ionization mass
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     Strege, Mark A.
     Lilly Corporate Center, Lilly Research Laboratories, Eli Lilly and Co.,
CS
     Indianapolis, IN, 46285, USA
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     Journal of Chromatography, B: Biomedical Sciences and Applications (1999),
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                    ARNOLD R, 1998, V18, P661, UNTERSUCHUNGEN ZUR BEEINFLUSSUNG
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                   HATHOWAY S, 1951, THE ATLAS FOR THE CLINICAL USE OF THE MMPI
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                   LEENDERS F, 1999, INVESTIGATION OF THE GRAMICIDIN S SYNTHETA
                   SE MULTIENZYME COMPLEX FOR BACILLUS BREVIS ATCC 999 PHD THES
                   IS TECHNICAL UNIVERSITY/RE
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         62843 FUNGI
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         41994 FUNGUS
            17 FUNGUSES
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(YEAST OR YEASTS)

E9

28

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                  CAPLUS
ТT
     Fungal metabolite screening: database of 474 mycotoxins and
     fungal metabolites for dereplication by standardised liquid
     chromatography-UV-mass spectrometry methodology
     Nielsen, Kristian Fog; Smedsgaard, Jorn
AU
CS
     BioCentrum-DTU, Mycology Group, Technical University of Denmark, Lyngby,
     DK-2800, Den.
     Journal of Chromatography, A (2003), 1002(1-2), 111-136
SO
     CODEN: JCRAEY; ISSN: 0021-9673
PB
     Elsevier Science B.V.
DT
     Journal
     English
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     ANSWER 2 OF 8 CAPLUS COPYRIGHT 2004 ACS on STN
     2002:557062 CAPLUS
AN
     137:259513
DN
     Direct surface analysis of fungal species by matrix-assisted
TI
     laser desorption/ionization mass spectrometry
AII
     Valentine, Nancy B.; Wahl, Jon H.; Kingsley, Mark T.; Wahl, Karen L.
CS
     Pacific Northwest National Laboratory, Richland, WA, 99352, USA
SO
     Rapid Communications in Mass Spectrometry (2002), 16(14), 1352-1357
     CODEN: RCMSEF; ISSN: 0951-4198
PR
     John Wiley & Sons Ltd.
DT
     Journal
T.A
     English
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     2002:456470 CAPLUS
AN
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     137:243340
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     Production of iturin A by Bacillus amyloliquefaciens suppressing
     Rhizoctonia solani
AU
     Yu, G. Y.; Sinclair, J. B.; Hartman, G. L.; Bertagnolli, B. L.
CS
     Department of Crop Sciences, University of Illinois at Urbana-Champaign,
     Urbana, IL, 61801-4709, USA
SO
     Soil Biology & Biochemistry (2002), 34(7), 955-963
     CODEN: SBIOAH; ISSN: 0038-0717
PB
     Elsevier Science Ltd.
DT
     Journal
LA
     English
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              ALL CITATIONS AVAILABLE IN THE RE FORMAT
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AN
     2002:298136 CAPLUS
DN
     137:19414
TT
     Integrated approach to explore the potential of marine microorganisms for
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Wagner-Dobler, Irene; Beil, Winfried; Lang, Siegmund; Meiners, Marinus;

Gesellschaft fur Biotechnologische Forschung, Mascheroder Weg 1,

the production of bioactive metabolites

Laatsch, Hartmut

Braunschweig, 38124, Germany

CS

- SO Advances in Biochemical Engineering/Biotechnology (2002), 74 (Tools and Applications of Biochemical Engineering Science), 207-238 CODEN: ABEBDZ; ISSN: 0724-6145
- PB Springer-Verlag
- DT Journal; General Review
- LA English
- RE.CNT 120 THERE ARE 120 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT
- L6 ANSWER 5 OF 8 CAPLUS COPYRIGHT 2004 ACS on STN
- AN 2002:195072 CAPLUS
- DN 137:334704
- TI Characterization of intact microorganisms by MALDI mass spectrometry
- AU Fenselau, Catherine; Demirev, Plamen A.
- CS Department of Chemistry and Biochemistry, University of Maryland, College Park, MD, 20742, USA
- SO Mass Spectrometry Reviews (2002), Volume Date 2001, 20(4), 157-171 CODEN: MSRVD3; ISSN: 0277-7037
- PB John Wiley & Sons, Inc.
- DT Journal; General Review
- LA English
- RE.CNT 113 THERE ARE 113 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT
- L6 ANSWER 6 OF 8 CAPLUS COPYRIGHT 2004 ACS on STN
- AN 2001:722112 CAPLUS
- DN 135:269520
- TI Characterization of protein biomarkers desorbed by MALDI from whole fungal cells
- AU Amiri-Eliasi, Bijan; Fenselau, Catherine
- CS Department of Chemistry and Biochemistry, University of Maryland, College Park, MD, 20742, USA
- SO Analytical Chemistry (2001), 73(21), 5228-5231 CODEN: ANCHAM; ISSN: 0003-2700
- PB American Chemical Society
- DT Journal
- LA English
- RE.CNT 31 THERE ARE 31 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT
- L6 ANSWER 7 OF 8 CAPLUS COPYRIGHT 2004 ACS on STN
- AN 2001:260581 CAPLUS
- TI Discrimination between bacterial spore types using time-of-flight mass spectrometry and matrix-free infrared laser desorption and ionization
- AU Ullom, J. N.; Frank, M.; Gard, E. E.; Horn, J. M.; Labov, S. E.; Langry, K.; Magnotta, F.; Stanion, K. A.; Hack, C. A.; Benner, W. H.
- CS Lawrence Livermore National Laboratory, Livermore, CA, 94550, USA
- SO Analytical Chemistry (2001), 73(10), 2331-2337 CODEN: ANCHAM; ISSN: 0003-2700
- PB American Chemical Society
- DT Journal
- LA English
- RE.CNT 23 THERE ARE 23 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT
- L6 ANSWER 8 OF 8 CAPLUS COPYRIGHT 2004 ACS on STN
- AN 2000:165884 CAPLUS
- DN 132:211777
- TI Detection and classification of individual airborne microparticles using laser ablation mass spectroscopy and multivariate analysis
- AU Parker, Eric P.; Trahan, Michael W.; Wagner, John S.; Rosenthal, Stephen E.; Whitten, William B.; Gieray, Rainer A.; Reilly, Peter T. A.; Lazar, Alexandru C.; Ramsey, J. Michael
- CS Department 15334: Laser and Computational Initiatives, Sandia National Laboratories, Albuquerque, NM, 87185-1188, USA

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SO Field Analytical Chemistry and Technology (2000), 4(1), 31-42 CODEN: FACTFR; ISSN: 1086-900X
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PB John Wiley & Sons, Inc.

DT Journal

LA English

RE.CNT 33 THERE ARE 33 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT

=> index bioscience FILE 'DRUGMONOG' ACCESS NOT AUTHORIZED

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ENTRY

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FULL ESTIMATED COST

62.60 62.81

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, AQUASCI, BIOBUSINESS, BIOCOMMERCE, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CANCERLIT, CAPLUS, CEABA-VTB, CEN, CIN, CONFSCI, CROPB, CROPU, DISSABS, DDFB, DDFU, DGENE, DRUGB, DRUGMONOG2, ...' ENTERED AT 15:17:11 ON 09 JAN 2004

68 FILES IN THE FILE LIST IN STNINDEX

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=> s (mass spectroscopy or mass spectrometry) and (fungi or fungus or fungal or pollen or yeast or mold)

- 393 FILE AGRICOLA
- 271 FILE ANABSTR
- 48 FILE AQUASCI
- 237 FILE BIOBUSINESS
- 3905 FILE BIOSIS
- 443 FILE BIOTECHABS
- 443 FILE BIOTECHDS
- 1566 FILE BIOTECHNO
- 925 FILE CABA
- 80 FILE CANCERLIT
- 2487 FILE CAPLUS
 - 85 FILE CEABA-VTB
 - 14 FILE CEN

17 FILES SEARCHED...

- 6 FILE CIN
- 9 FILE CONFSCI
- 23 FILE CROPB
- 23 FILE CROPU
- 192 FILE DISSABS
- 49 FILE DDFB
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- 749 FILE DGENE
- 49 FILE DRUGB
- 80 FILE DRUGU 29 FILES SEARCHED...
 - 36 FILE EMBAL
 - 2737 FILE EMBASE
 - 1000 FILE ESBIOBASE
 - 439 FILE FEDRIP
 - 188 FILE FROSTI
 - 112 FILE FSTA
 - 14 FILE GENBANK
 - 9 FILE HEALSAFE
 - 124 FILE IFIPAT
 - 539 FILE JICST-EPLUS
 - 9 FILE KOSMET
 - 795 FILE LIFESCI
 - 1288 FILE MEDLINE

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FILE IFIPAT

FILE JICST-EPLUS

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 - 5 FILE TOXCENTER
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- 19 FILES HAVE ONE OR MORE ANSWERS, 68 FILES SEARCHED IN STNINDEX
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F2		12	CAPLUS
F3		11	SCISEARCH
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F19		1	PHIN

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PΒ

John Wiley & Sons Ltd.

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The intact fungal spores of several strains of 4 Aspergillus
AB
     species, Aspergillus flavus, A. oryzae, A. parasiticus, and A. sojae, were
     directly analyzed by matrix-assisted laser desorption/ionization (
     MALDI) time-of-flight mass
     spectrometry. Very simple MALDI mass spectra are obtained by
     directly mixing spores with a matrix such as .alpha.-cyano-4-
     hydroxycinnamic acid or sinapinic acid. The mass spectra are obtained
     from the ablation of cell walls of spores owing to the acidity of the
     matrix soln. The MALDI results show that aflatoxigenic strains and
     non-aflatoxigenic strains have different mass peak profiles.
     Furthermore, the MALDI results of non-aflatoxigenic A. flavus and A.
     parasiticus spores resemble those of the closely related A. oryzae and A.
     sojae spores, resp.
RE.CNT 21
              THERE ARE 21 CITED REFERENCES AVAILABLE FOR THIS RECORD
              ALL CITATIONS AVAILABLE IN THE RE FORMAT
     ANSWER 2 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 2
L12
AN
     2001:15096 CAPLUS
DN
     134:246820
TT
     Identification of an N-(hydroxysulfonyl)oxy metabolite using in vitro
     microorganism screening, high-resolution and tandem electrospray
     ionization mass spectrometry
ΑU
     Pilard, Serge; Caradec, Fabrice; Jackson, Peter; Luijten, Wim
CS
     UPJV, Laboratoire des Glucides, Amiens, 80039, Fr.
SO
     Rapid Communications in Mass Spectrometry (2000), 14(24),
     2362-2366
     CODEN: RCMSEF; ISSN: 0951-4198
PB
     John Wiley & Sons Ltd.
DT
     Journal
LA
     English
     Preliminary metabolic profiling of a drug under pre-clin.
AB
     development (S 19812) revealed the presence of a minor unknown metabolite
     with a pos. ion electrospray ionization (ESI) mass spectrum identical to
     that of the unchanged compd. Since the low concn. of the compd. did not
     allow any addnl. expts., preparative bioconversion using fungi
     was used to obtain a substantial amt. of the mol. Neq. ion ESI
     -MS and tandem mass spectrometry (MS/MS) in
     combination with accurate mass measurements obtained on a
     quadrupole/time-of-flight instrument (Q-TOF) led to the pos.
     identification of a hydroxylamide sulfoconjugated metabolite.
              THERE ARE 9 CITED REFERENCES AVAILABLE FOR THIS RECORD
RE.CNT 9
              ALL CITATIONS AVAILABLE IN THE RE FORMAT
L12
     ANSWER 3 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 3
ΑN
     2001:65927 CAPLUS
DN
     134:211383
     Design and microstructuring of PDMS surfaces for improved marine
TI
     biofouling resistance
AU
     Petronis, Sarunas; Berntsson, Kent; Gold, Julie; Gatenholm, Paul
CS
     Department of Applied Physics, Chalmers University of Technology,
     Goeteborg, SE-412 96, Swed.
SO
     Journal of Biomaterials Science, Polymer Edition (2000), 11(10),
     1051-1072
     CODEN: JBSEEA; ISSN: 0920-5063
PB
     VSP BV
DT
     Journal
LA
     English
AB
     In this study room temp. vulcanized (RTV) silicone surfaces with designed
     surface microstructure and well-defined surface chem. were prepd. Their
     resistance to marine macrofouling by barnacles Balanus improvisus was
     tested in field expts. for deducing optimal surface topog. dimensions
     together with a better understanding of macrofouling mechanisms.
     Polydimethylsiloxane (PDMS) surfaces were microstructured by casting the
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DT

LΑ

Journal English

PDMS pre-polymer on microfabricated molds. The master molds were made by utilizing photolithog, and anisotropic etching of monocryst, silicon wafers. Several iterative casting steps of PDMS and epoxy were used to produce large quantities of microstructured PDMS samples for field studies. The microstructured PDMS surface consisted of arrays of pyramids or riblets creating a surface arithmetic mean roughness ranging from 5 to 17 .mu.m for different microstructure sizes and geometries, as detd. by SEM. Chemophys, properties of the microstructured films were investigated by electron spectroscopy for chem. anal., time-of-flight secondary ion mass

spectroscopy and dynamic contact angle measurements. Films were chem. homogeneous down to the submicron level. Hydrophobicity and contact angle hysteresis increased with increased surface roughness. Field tests on the west coast of Sweden revealed that the microstructure contg. the largest riblets (profile height 69 .mu.m) reduced the settling of barnacles by 67%, whereas the smallest pyramids had no significant influence on settling compared to smooth PDMS surfaces. The effect of dimensions and geometry of the surface microstructures on the B. improvisus larvae settling is discussed.

RE.CNT 36 THERE ARE 36 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT

- L12 ANSWER 4 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 4
- AN 2000:159323 CAPLUS
- DN 132:319392
- TI Characterization of **fungal** spores by laser desorption/ionization time-of-flight mass spectrometry
- AU Welham, K. J.; Domin, M. A.; Johnson, K.; Jones, L.; Ashton, D. S.
- CS ULIRS Mass Spectrometry Laboratory, Department of Pharmaceutical and Biological Chemistry, The School of Pharmacy, University of London, London, WC1N 1AX, UK
- SO Rapid Communications in Mass Spectrometry (2000), 14(5), 307-310 CODEN: RCMSEF; ISSN: 0951-4198
- PB John Wiley & Sons Ltd.
- DT Journal
- LA English
- AB A considerable vol. of research has now been completed on the application of matrix-assisted laser desorption/ionization mass spectrometry (MALDI-MS) to the anal. of

bacteria; however, to date no definitive studies have been made using this technique on **fungi**. Preliminary studies on the application of the MALDI-MS methodol., previously developed for the anal. of bacteria, to the anal. of intact **fungal** spores are described here.

MALDI-MS and electrospray mass

spectrometry enable the high mol. wt. anal. of proteins, glycoproteins, oligosaccharides and oligonucleotides. Using MALDI-MS with bacteria has demonstrated the ability to produce "fingerprints" of the intact cells with the ions obsd. being assocd. with the proteinaceous components of the cell wall. This paper reports the adaptation of this technique to the direct anal. of fungal cells. The high percentage of carbohydrate in the fungal cell wall indicates that the ions obsd. in the mass spectrometric expts. may be of carbohydrate origin. Penicillium spp., Scytalidium dimidiatum and Trichophyton rubrum have been studied in this preliminary investigation and all show individually distinctive spectra which would appear to provide a profile of the cellular material with discrete peaks being obsd. over the mass range 2 to 13 kDa. The spectra obtained are reproducible within the method used but, as shown in our previous studies on bacteria, washing may selectively release components from the fungal cell wall.

RE.CNT 3 THERE ARE 3 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT

L12 ANSWER 5 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 5 AN 2000:323975 CAPLUS

- DN 133:86552
- TI New sequences and new fungal producers of peptaibol antibiotics antiamoebins
- AU Jaworski, Andreas; Bruckner, Hans
- CS Department of Food Sciences, Institute of Nutritional Science, University of Giessen, Giessen, 35390, Germany
- SO Journal of Peptide Science (2000), 6(4), 149-167 CODEN: JPSIEI; ISSN: 1075-2617
- PB John Wiley & Sons Ltd.
- DT Journal
- LA English
- Mixts. of the microheterogeneous 16-mer peptaibol antibiotics called AB antiamoebins [AAM] have been isolated from the culture broths of strains of the filamentous fungi Stilbella erythrocephala ATCC 28144, Stilbella fimetaria CBS 548.84 and Gliocladium catenulatum CBS 511.66. Sequences were detd. using online HPLC together with pos.- and neg.-ion electrospray ionization mass spectrometry. Some characteristic features are recognized in the mass spectrometric fragmentation pattern of AAM. From a sample originally used for sequencing AAM (from Hindustan Antibiotics, Ltd., Pimpri, Poona-411018, India), and a sample of AAM com. available (from Sigma Chems., St. Louis, MO, USA) HPLC elution profiles and sequences were assigned. Further, sequences of AAM previously isolated from Emericellopsis synnematicola CBS 176.60 and Emericellopsis salmosynnemata CBS 382.62 were detd. The peptide designated AAM I was the most abundant in all isolates and its structure could be confirmed. AAM II was detectable as a minor component (1.9%) only in the original sample of AAM, but not in the other The structures of AAM III, IV and V, which had previously been isolates. partly assigned, were definitely established, and the new sequences AAM VI-XVI were elucidated. AAM showing Phel/Leu1 or Phel/Val1 exchange, resp., are produced in amts. only by S. erythrocephala. Sequences, HPLC elution profiles ("fingerprints") and relative amts. of peptides of all isolates were correlated.
- RE.CNT 51 THERE ARE 51 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT
- L12 ANSWER 6 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 6
- AN 2000:258180 CAPLUS
- DN 133:14253
- TI Matrix-assisted laser-desorption/ionization time-offlight mass spectrometry and its application to the analysis of fungal spores
- AU Welham, K. J.; Domin, M. A.; Johnson, K.; Jones, L.; Ashton, D. S.
- CS ULIRS Mass Spectrometry Laboratory, Department of Pharmaceutical and Biological Chemistry, The School of Pharmacy, University of London, London, WC1N 1AX, UK
- SO Pharmacy and Pharmacology Communications (2000), 6(3), 107-111 CODEN: PPCOFN; ISSN: 1460-8081
- PB Royal Pharmaceutical Society of Great Britain
- DT Journal
- LA English AB Although much research has been completed on the application of matrix-assisted laser-desorption/ionization mass spectrometry (MALDI-MS) to the anal. of bacteria, no definitive studies have yet been performed on the anal. of fungi Preliminary studies on the application of the MALDI-MS methodol., previously developed for the anal. of bacteria, to the anal. of intact fungal spores are described here. MALDI-MS and electrospray mass spectrometry enable the anal. of high mol.-wt. proteins, glycoproteins, oligosaccharides and oligonucleotides. Using MALDI-MS with bacteria has enabled the prodn. of "fingerprints" of the intact cells; the ions obsd. are assocd. with the proteinaceous components of the cell wall. This study reports the adaptation of this technique to the direct anal. of fungal

cells. Because of the large amt. of carbohydrate in the fungal

cell wall, the ions obsd. in the mass spectrometric expts. might be of carbohydrate origin. Penicillium spp., Scytalidium dimidiatum and Trichophyton rubrum have been studied in this preliminary investigation and all furnish individually distinctive spectra which seem to provide a profile of the cellular material with discrete peaks being obsd. over the mass range 2 to 13 kDa. The spectra obtained are reproducible within the method used but, as shown in our previous studies on bacteria, washing might selectively release components from the fungal cell wall.

RE.CNT 3 THERE ARE 3 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT

- L12 ANSWER 7 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 7
- AN 1999:560853 CAPLUS
- DN 132:60756
- TI Electrospray ionization tandem mass spectrometry (ESI-MS/MS) analysis of the lipid molecular species composition of yeast subcellular membranes reveals acyl chain-based sorting/remodeling of distinct molecular species en route to the plasma membrane
- AU Schneiter, Roger; Brugger, Britta; Sandhoff, Roger; Zellnig, Gunther; Leber, Andrea; Lampl, Manfred; Athenstaedt, Karin; Hrastnik, Claudia; Eder, Sandra; Daum, Gunther; Paltauf, Fritz; Wieland, Felix T.; Kohlwein, Sepp D.
- CS Spezialforschungsbereich Biomembrane Research Center, Institut fur Biochemie und Lebensmittelchemie, Technische Universitat Graz, Graz, A-8010, Austria
- SO Journal of Cell Biology (1999), 146(4), 741-754 CODEN: JCLBA3; ISSN: 0021-9525
- PB Rockefeller University Press
- DT Journal
- LA English
- AB Nano-electrospray ionization tandem mass spectrometry (nano-ESI-MS/MS) was employed to det. qual. differences in the lipid mol. species compn. of a comprehensive set of organellar membranes, isolated from a single culture of Saccharomyces cerevisiae cells. Marked differences in the acyl chain compn. of biosynthetically related phospholipid classes were obsd. Acyl chain satn. was lowest in phosphatidylcholine (PC, 15.4%) and phosphatidylethanolamine (PE, 16.2%), followed by phosphatidylserine (PS, 29.4%), and highest in phosphatidylinositol (PI, 53.1%). The lipid mol. species profiles of the various membranes were generally similar, with a deviation from a calcd. av. profile of approx. .+-.20%. Nevertheless, clear distinctions between the mol. species profiles of different membranes were obsd., suggesting that lipid sorting mechanisms were operating at the level of individual mol. species to maintain the specific lipid compn. of a given membrane. Most notably, the plasma membrane was enriched in satd. species of PS and PE. The nature of the sorting mechanism that dets. the lipid compn. of the plasma membrane was investigated further. The accumulation of mono-unsatd. species of PS at the expense of di-unsatd. species in the plasma membrane of wild-type cells was reversed in elo3.DELTA. mutant cells, which synthesize C24 fatty acid-substituted sphingolipids instead of the normal C26 fatty acid-substituted species. This observation suggests that acyl chain-based sorting and/or remodeling mechanisms are operating to maintain the specific lipid mol. species compn. of the yeast plasma membrane.
- RE.CNT 73 THERE ARE 73 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT
- L12 ANSWER 8 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 8
- AN 1999:129007 CAPLUS
- DN 130:301822
- TI Online nonaqueous capillary electrophoresis and electrospray mass spectrometry of tricyclic antidepressants and metabolic **profiling** of amitriptyline by Cunninghamella elegans

- Liu, Chun-Sheng; Li, Xing-Fang; Pinto, Devanand; Hansen, Eugene B., Jr.; AU Cerniglia, Carl E.; Dovichi, Norman J.
- CS Department Chemistry, University Alberta, Edmonton, AB, Can.
- Electrophoresis (1998), 19(18), 3183-3189 SO CODEN: ELCTDN; ISSN: 0173-0835
- PBWiley-VCH Verlag GmbH
- DTJournal
- LA English
- An online nonaq. capillary electrophoresis-electrospray AB mass spectrometry (ESI-MS) was developed using a com. ion spray interface. The nonaq. capillary electrophoresis ESI-MS system was used to profile tricyclic antidepressants of similar structures and mass-to-charge ratios. The authors found that pure MeOH can be used as a sheath liq. to obtain stable ion spray from nonaq. capillary electrophoresis. The flow rate of the coaxial nebulizing gas affected baseline signals, sepn. efficiency, and migration times. Other nonag. capillary electrophoresis operating conditions and electrospray parameters were optimized for enhanced baseline sepn. and high sensitivity detection. The effect of sample stacking on sepn. and detection was The calcd. detection limits were 3 pg injected onto the evaluated. capillary. ESI mass spectra of tricyclic antidepressants from a single quadrupole MS were obtained and elucidated. The information was used to propose fragmentation pathways of the tricyclic antidepressants. method was also used to analyze the metabolites of amitriptyline produced by the fungus C. elegans. 16 Metabolites were detected and most of them were tentatively identified as demethylated and/or hydroxylated, and/or N-oxidized products.
- RE.CNT 22 THERE ARE 22 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT
- ANSWER 9 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 9 L12
- $\mathbf{A}\mathbf{N}$ 1997:403133 CAPLUS
- DN 127:94242
- HPLC-MS/MS Profiling of Tryptophan-Derived Alkaloids in Food: TT Identification of Tetrahydro-.beta.-carbolinedicarboxylic Acids
- ΑU Gutsche, B.; Herderich, M.
- Lehrstuhl fuer Lebensmittelchemie, Universitaet Wuerzburg, Wuerzburg, CS 97074, Germany
- SO Journal of Agricultural and Food Chemistry (1997), 45(7), 2458-2462 CODEN: JAFCAU; ISSN: 0021-8561
- PB American Chemical Society
- DTJournal
- LA English
- AB A method for selective detection of 1,2,3,4-tetrahydro-.beta.carbolinecarboxylic acids (THCCs) was developed based on electrospray ionization-tandem mass spectrometry coupled to liq. chromatog. (HPLC-ESI-MS/MS). Low-energy collision-induced dissocn. (CID) led to characteristic fragment ions due to neutral loss of 73 amu. Subsequently, const. neutral loss scanning was used for substructure specific screening of THCCs in food samples. Detection limits for HPLC-ESI-MS/MS anal. of THCCs applying neutral loss expts. were established at 100 ng/mL (ca. 2.5 pmol on column). Application of this MS/MS method enabled us to detect THCC derivs. derived from Pictet-Spengler condensation of tryptophan with .alpha.-oxo acids. Subsequently, diastereomeric 1,2,3,4-tetrahydro-.beta.-carboline-1,3dicarboxylic acid 3a/b, 1-methyl-1,2,3,4-tetrahydro-.beta.-carboline-1,3dicarboxylic acid 4a/b, and 1-(2'-carboxyethyl)-1,2,3,4-tetrahydro-.beta.carboline-3-carboxylic acid 5a/b were identified in alc. beverages, seasoning sauces, yeast ext., and fruit products for the 1st time. Most food samples under study contained 3a/b and 4a/b in significant amts. 5A/b was identified in soy sauce, worcestershire sauce, seasoning sauce, and yeast ext. Due to the excellent selectivity of tandem mass spectrometry coeluting tetrahydro-.beta.carboline derivs. could be identified unequivocally by HPLC-ESI-MS/MS.

- L12 ANSWER 10 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 10
- AN 1997:335519 CAPLUS
- DN 127:62808
- TI Identification of two-dimensional gel electrophoresis resolved **yeast** proteins by **matrix-assisted** laser desorption ionization **mass spectrometry**
- AU Larsson, Thomas; Norbeck, Joakim; Karlsson, Hasse; Karlsson, Karl-Anders; Blomberg, Anders
- CS Medical Biochemistry, Goteborg University, Goteborg, Swed.
- SO Electrophoresis (1997), 18(3-4), 418-423 CODEN: ELCTDN; ISSN: 0173-0835
- PB VCH
- DT Journal
- LA English
- Protein ext. from yeast cells growing exponentially in saline AB medium was sepd. by two-dimensional PAGE (2-D PAGE), with the sepn. in the first dimension on a wide range immobilized pH (3-10) gradient. From one preparative 2-D gel a no. of previously identified proteins were used as test material for our initial matrix-assisted laser desorption ionization mass spectrometry (MALDI -MS) efforts on large scale rapid protein spot identification. Sample prepn. via in-gel trypsin digestion was slightly modified to be compatible to MS anal., and via this modified procedure MS generated peptide mass profiles could, in most cases with good precision, identify the protein in question. Preferential ionization was tested on a yeast aldehyde dehydrogenase (ALD7), and it was shown that the ionization of some peptides was clearly suppressed by the presence of others. Roughly 50% of the obsd. peptide masses was found by the search routines in the database, and the mass measurement accuracy of the peptides was within 0.5 Da. Silver-stained gels could be used with good results for the generation of peptides to be analyzed by MALDI-MS. For one of the 2-D resolved proteins, glycerol 3-phosphatase (GPP1), the post-source decay (PSD) spectrum proved crucial in identification.
- L12 ANSWER 11 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN DUPLICATE 11
- AN 1996:330368 CAPLUS
- DN 125:52639
- TI Using direct electrospray mass spectrometry in taxonomy and secondary metabolite profiling of crude fungal extracts
- AU Smedsgaard, Joern; Frisvad, Jens C.
- CS Department Biotechnology, Technical University Denmark, Lyngby, 2800, Den.
- SO Journal of Microbiological Methods (1996), 25(1), 5-17 CODEN: JMIMDQ; ISSN: 0167-7012
- PB Elsevier
- DT Journal
- LA English
- AB Important information about sample compn. can be obtained within a few minutes by injecting a complex mixt. like a crude ext. of a fungal culture prepd. for std. HPLC anal. directly into an electrospray mass spectrometer using an FIA-ESMS type of setup. The limited fragmentation and high sensitivity of ESMS were used in this study to provide mass profiles from ethylacetate/methanol/chloroform exts. from cultures of 10 of the most common Penicillium species assocd. with stored cereals. The anal. parameters were optimized to reduce fragmentations and reactions in ESMS; hence only the protonated or sodiated ions were obsd. for most The anal. demonstrated that ions corresponding to the protonated mol. ions (M + H+) from most of the known secondary metabolites and mycotoxins produced by these species could be obsd. in the ESMS spectra. A no. of other distinct species-specific ion were obsd. as well. The 10 different species could be discriminated either by ions corresponding to known or unknown metabolites. By creating a database of mass spectra obtained from anal. of different species using the facility included in std. MS software, it was possible to use a simple library search to

identify most of the species included in this study on the basis of their mass spectra.

- L12 ANSWER 12 OF 19 CAPLUS COPYRIGHT 2004 ACS on STN
- AN 1997:372822 CAPLUS
- DN 127:62826
- TI Identification and detection of bacteria: electrospray MS-MS versus derivatization/GC-MS
- AU Fox, Alvin; Black, Gavin; Fox, Karen; Wunschel, David
- CS Dept. Microbiol. Immunol., USC School Medicine, Columbia, SC, 29208, USA
- Proceedings of the ERDEC Scientific Conference on Chemical and Biological Defense Research, Aberdeen Proving Ground, Md., Nov. 15-18, 1994 (1996), Meeting Date 1994, 39-44. Editor(s): Berg, Dorothy A. Publisher: National Technical Information Service, Springfield, Va. CODEN: 64NAAX
- DT Conference
- LA English
- AB Identification of chem. markers and demonstration of their taxonomic utility is essential in conjunction with development of a real-time biodetection strategy. Pyrolysis MS-MS is an attractive technique in terms of rapidity of anal. Unfortunately, many marker compds. are destroyed by the drastic depolymn./volatilization conditions. A gentler alternative is electrospray ionization tandem mass spectrometry (ESI MS-MS) which generates simple parent ion spectra and readily interpretable product spectra. This lab. has developed a scheme for identification of Bacillus anthracis, a key biol. agent, using carbohydrate profiling of spores and vegetative cells identified as alditol acetate derivs. by GC-MS and 16S/23S interspace region PCR products. Further characterization of these chem. markers is in progress. Using ESI-MS-MS, we have used muramic acid as a general marker for bacteria allowing their ready differentiation from fungi. Sample prepn. is minimal; hydrolysis in acid is followed by extn. with an org. base (or barium carbonate neutralization). logical extension of this work would include anal. of monomers and polymers, (naturally present in microbial cells) without prior depolymn. We have demonstrated that dipicolinic acid (DPA, a marker for spores) on ESI generates a mol. ion as a parent, M-H- (m/z 166) and characteristic daughter ions in MS-MS mode $[m/z\ 122\ (loss\ of\ CO2)\ and\ m/z$ 78 (loss of 2CO2)]. In contrast, pyrolysis destroys DPA producing pyridine by decarboxylation (seen as m/z 79) in the parent spectrum. We aim to demonstrate that ESI MS-MS anal. for DPA can specifically differentiate spores from vegetative cells. Mols. with more general potential in biodetection include nucleotides coded by the rRNA operon. These mols. are universally found in high concn. in bacterial cells but can also be amplified by PCR. Encouraging preliminary results have been obtained with MS-MS anal. of synthetic oligonucleotides. Obsd. and calcd. m/z values are in close agreement. There is considerable potential for ESI MS-MS in rapid biodetection of monomers, oligomers and polymers.
- L12 ANSWER 13 OF 19 SCISEARCH COPYRIGHT 2004 THOMSON ISI on STN
- AN 94:774977 SCISEARCH
- GA The Genuine Article (R) Number: PU984
- TI ELECTROSPRAY TANDEM MASS-SPECTROMETRY FOR ANALYSIS OF NATIVE MURAMIC ACID IN WHOLE BACTERIAL-CELL HYDROLYSATES
- AU BLACK G E; FOX A (Reprint); FOX K; SNYDER A P; SMITH P B W
- CS UNIV S CAROLINA, SCH MED, COLUMBIA, SC, 29208 (Reprint); UNIV S CAROLINA, SCH MED, COLUMBIA, SC, 29208; USA CHEM RES, CTR DEV & ENGN, ABERDEEN PROVING GROUND, MD, 21010; GEOCENTERS INC, GUNPOWDER BRANCH, ABERDEEN PROVING GROUND, MD, 21010
- CYA USA
- SO ANALYTICAL CHEMISTRY, (01 DEC 1994) Vol. 66, No. 23, pp. 4171-4176.
 ISSN: 0003-2700.
- DT Article; Journal

FS PHYS; LIFE

LA ENGLISH

AΒ

REC Reference Count: 19

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

Muramic acid is an amino sugar found in eubacterial cell walls and not elsewhere in nature. This study explored the use of electrospray tandem mass spectrometry (ESI MS/MS) in analysis of underivatized muramic acid in bacterial hydrolysates. Fungal hydrolysates were used as negative controls. The only processing used was hydrolysis in sulfuric acid followed by extraction with an organic base (N,N-dioctylmethylamine) to remove the acid prior to ESI MS/MS analysis. Compared with pure muramic acid, bacterial hydrolysates produced more complex ESI mass spectra, such that the protonated molecular ion at m/z 252 was barely detectable. In contrast, product ion spectra of m/z 252 were identical among pure muramic acid, Gram positive bacteria, and Gram negative bacteria. However, no characteristic product ion spectrum was manifested from m/z 252 in fungal samples. This allowed ready, visual differentiation of bacteria and fungi. Multiple reaction monitoring (MRM) following muramic acid fragmentations (m/z 252 --> 144 and m/z 252 --> 126) increased sensitivity and allowed quantitative differentiation when compared with the MRM of the internal standard N-methyl-D-glucamine (m/z 196 --> 44). ESI MS/MS required minimal sample preparation and allowed rapid sample throughput for analysis of muramic acid in whole bacterial cell hydrolysates.

L12 ANSWER 14 OF 19 PROMT COPYRIGHT 2004 Gale Group on STN

AN 2000:57338 PROMT

TI Manufacturers and Suppliers. (Alphabetical list of companies)

SO Lasers & Optronics, (Nov 1999) Vol. 18, No. 11, pp. S8. ISSN: 0892-9947.

Cahners Publishing Company

DT Newsletter

LA English

WC 71777

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AB A

PB

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AN 2000:2470 PROMT

TI EUROPEAN PATENT DISCLOSURES PRIVATE.

SO BIOWORLD Today, (29 Dec 1999) Vol. 10, No. 247.

PB American Health Consultants, Inc.

DT Newsletter

LA English

WC 2624

FULL TEXT IS AVAILABLE IN THE ALL FORMAT

AB Acacia Biosciences WO 99/58720 Gene expression **profiling**Kirkland, Wash. Methods, systems, and instrumentation for quantifying the relatedness of gene expression **profiles**.

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Subscription: \$1350.00 per year. Published daily (5 times a week). Box 740021, Atlanta, GA 30374.

L12 ANSWER 16 OF 19 PROMT COPYRIGHT 2004 Gale Group on STN

AN 94:582623 PROMT

TI Analytical chemistry. (advancements in separation science)

AU Russell, David A.

- SO Chemistry and Industry, (3 Oct 1994) No. 19, pp. 783(2). ISSN: ISSN: 0009-3068.
- PB Society of Chemical Industry
- DT Newsletter
- LA English
- WC 1073
 - *FULL TEXT IS AVAILABLE IN THE ALL FORMAT*
- AB Over the last review period there has been a number of reported developments in the area of separation science. The separation of large DNA fragments has been achieved using pulsed field capillary zone electrophoresis (CZE) (J Sudor and M V Novotny, Anal. Chem., 1994, 66, 2446). The CZE technique, using an entangled polyacrylamide solution, was applied to large DNA samples under pulsed-field conditions. Highly efficient separations were obtained using biased sinusoidal field and field-inversion pulsing regimes. [Lambda]DNA standards (8.348.5kb) and 48.5kb-1Mb [Lambda]DNA concatamers clearly demonstrated an improved separation time of about 10-50 times over conventional slab-gel techniques. The authors also suggested that the CZE method appeared more sensitive and amenable to component quantification and method automation.
- L12 ANSWER 17 OF 19 BIOBUSINESS COPYRIGHT 2004 BIOSIS on STN
- AN 97:66571 BIOBUSINESS
- DN 0924106
- TI HPLC-MS-MS **profiling** of tryptophan-derived alkaloids in food: Identification of tetrahydro-beta-carbolinedicarboxylic acids.
- AU Gutsche B; Herderich M
- CS Lehrstuhl Lebensmittelchem., Univ. Wuerzburg, Am Hubland, 97074 Wuerzburg, Germany.
- SO Journal of Agricultural and Food Chemistry, (1997) Vol.45, No.7, p.2458-2462.
 ISSN: 0021-8561.
- DT ARTICLE
- FS NONUNIQUE
- LA English
- AB A method for selective detection of 1,2,3,4-tetrahydro-betacarbolinecarboxylic acids (THCCs) was developed based on electrospray ionization-tandem mass spectrometry coupled to liquid chromatography (HPLC-ESI-MS/MS). Low-energy collision-induced dissociation (CID) led to characteristic fragment ions due to neutral loss of 73 amu. Subsequently, constant neutral loss scanning was used for substructure specific screening of THCCs in food samples. Detection limits for HPLC-ESI-MS/MS analysis of THCCs applying neutral loss experiments were established at 100 ng mL-1 (ca. 2.5 pmol on column). Application of this MS/MS method enabled us to detect THCC derivatives derived from Pictet-Spengler condensation of tryptophan with alpha-oxo acids. Subsequently, diastereomeric 1,2,3,4-tetrahydro-betacarboline-1,3-dicarboxylic acid 3a/b, 1-methyl-1,2,3,4-tetrahydro-betacarboline-1,3-dicarboxylic acid 4a/b, and 1-(2'-carboxyethyl)-1,2,3,4tetrahydro-beta-carboline-3-carboxylic acid 5a/b were identified in alcoholic beverages, seasoning sauces, yeast extract, and fruit products for the first time. Most food samples under study contained 3a/b and 4a/b in significant amounts. 5a/b was identified in soy sauce, worcestershire sauce, seasoning sauce, and yeast extract. Due to the excellent selectivity of tandem mass spectrometry coeluting tetrahydro-beta-carboline derivatives could be identified unequivocally by HPLC-ESI-MS/MS.
- L12 ANSWER 18 OF 19 DISSABS COPYRIGHT (C) 2004 ProQuest Information and Learning Company; All Rights Reserved on STN
- AN 2001:58769 DISSABS Order Number: AAIMQ57123
- TI Characterization of protein glycoforms in cellobiohydrolases and endoglucanases from Trichoderma reesei RUT-C30 and mutant strains using capillary isoelectric focusing and mass spectrometry
- AU Hui, Joseph P. M. [M.Sc.]; Roy, Rene [adviser]; Thibault, Pierre [adviser]
- CS University of Ottawa (Canada) (0918)

SO Masters Abstracts International, (2000) Vol. 39, No. 4, p. 1162. Order No.: AAIMQ57123. 142 pages. ISBN: 0-612-57123-8.

Dissertation

FS MAI

DT

AB

LA English

<italic>Trichoderma reesei</italic> is a filamentous fungus heavily used in the biotechnology industry due to its efficient secretion of cellulases. The enzymatic system of <italic>T. reesei</italic> consists primarily of four glycoproteins referred to as cellobiohydrolases (CBH I, CBH II) and endoglucanases (EG I, EG II). They exhibit microheterogeneity both in the N- and O-linked glycans. This thesis focuses on the method development to characterize the glycosylation profile and post-translational modifications present in these glycoproteins. Crude cellulase fermentation extracts RUT-C30 and its two mutant strains Iogen-M4, Iogen-B13 were initially analyzed by capillary isoelectric focusing (CIEF) to determine the cellulase composition in them. The major cellulase CBH I was purified from each strain and electrospray mass spectrometry (ESMS) was used to reveal the extent of overall glycosylation. To characterize the N-linked glycans and their attachment sites, CBH I from these strains were subjected to tryptic digest with and without PNGase F incubations followed by mass spectrometric detection. The O-linked glycans were released chemically by hydrazinolysis and were analyzed by high performance anion-exchange chromatography with pulsed amperometric detection (HPAEC-PAD). The majority of O-linked glycans was di- and tri-saccharides. Two unusual posttranslational modification from strain RUT-C30 were observed: (1) both high mannose (predominantly Man₈GlcNAc₂) and single GlcNAc in putative N-linked sites and (2) mannosylphosphorylation in a O-linked di-saccharide. Heterogeneity in putative N-linked sites was found consistently in CBH II, EG I and EG II from RUT-C30, however, no mannosylphosphorylation was observed at least on the proteins in the purified fractions. These results have led to the proposal of endogenous endoglycosidase H as well as mannosylphosphorylation activities possibly induced during fermentation.

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AN 1998:8050 PHIN

DN B00577672

DED 1 Apr 1998

TI Pharmacogenomic Strategies: Biotech's New David and Goliath Challenge

SO Bioventure-View (1998) No. 1304 p1

DT Newsletter

FS FULL

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